plication No.: <u>09/530,663</u>

## NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

PE VOY		<ol> <li>This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.</li> </ol>
RADEMARY		<ol> <li>This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).</li> </ol>
MAG		3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
	x	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
		5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
		6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
		7. Other:
	Apı	olicant Must Provide:
	x	An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
	x	An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
	X	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
	For	questions regarding compliance to these requirements, please contact:
		Rules Interpretation, call (703) 308-4216
		CRF Submission Help, call (703) 308-4212 entIn Software Program Support (SIRA)
		Technical Assistance 703-287-0200

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A



#9 / 1635 2-23-d635

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/530,663

DATE: 02/16/2001 TIME: 16:36:30

Input Set : A:\B0045591.txt

Output Set: N:\CRF3\02162001\1530663.raw

Does Not Comply Corrected Diskette Needed

## ERRORED SEQUENCES

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i is invalid in the sequence itself-use"n,"

sequence itself-use"n,"

and Aplain

and Aplain

28 in 62207-62237

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      563 <211> LENGTH: 31
      564 <212> TYPE: DNA
      565 <213> ORGANISM: Populus tremuloides Michx. (aspen)
567 <400> SEQUENCE: 9
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570 <210> SEQ ID NO: 10
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      572 <212> TYPE: DNA
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      575 <400> SEQUENCE: 10
E--> 576 ttggatccgt (igcicarcar gt/gaygg Same LWW
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      579 <211> LENGTH: 27
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      583 <400> SEQUENCE: 11
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                                                                                              27
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      629 <211> LENGTH: 11
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      633 <400> SEQUENCE: 17
                                                                                               11
      634
           tctcaccaac c
E--> 640(1)
E--> 643(2) delete at end of file
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/530,663

DATE: 02/16/2001 TIME: 16:36:31

Input Set : A:\B0045591.txt

Output Set: N:\CRF3\02162001\I530663.raw

L:568 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6 L:576 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:3 L:584 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:640 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:11 SEQ:17

M:254 Repeated in SeqNo=17